



X11600.ST25.txt  
SEQUENCE LISTING

<110> Becker, Gerald  
Hale, John  
Heath, William  
Johnstone, Edward  
Little, Sheila  
Tu, Yuan  
Yeh, Wu-Kuang  
Yin, Tinggui

<120> Amyloid Precursor Protein Protease and Related Nucleic Acid  
Compounds

<130> X-11600

<160> 5

<170> PatentIn version 3.1

<210> 1

<211> 1683

<212> DNA

<213> Homo sapiens

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<222> (25)..(1629)

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<221> misc\_feature

<222> (25)..(1629)

X11600.ST25.txt

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atg ggt gtg cat gtg ggt gca gcc ctg gga gca ctg tgg ttc tgc ctc      99
Met Gly Val His Val Gly Ala Ala Leu Gly Ala Leu Trp Phe Cys Leu
10                15                20                25

aca gga gcc ctg gag gtc cag gtc cct gaa gac cca gtg gtg gca ctg      147
Thr Gly Ala Leu Glu Val Gln Val Pro Glu Asp Pro Val Val Ala Leu
30                35                40

gtg ggc acc gat gcc acc ctg tgc tgc tcc ttc tcc cct gag cct ggc      195
Val Gly Thr Asp Ala Thr Leu Cys Cys Ser Phe Ser Pro Glu Pro Gly
45                50                55

ttc agc ctg gca cag ctc aac ctc atc tgg cag ctg aca gat acc aaa      243
Phe Ser Leu Ala Gln Leu Asn Leu Ile Trp Gln Leu Thr Asp Thr Lys
60                65                70

cag ctg gtg cac agc ttt gct gag ggc cag gac cag ggc agc gcc tat      291
Gln Leu Val His Ser Phe Ala Glu Gly Gln Asp Gln Gly Ser Ala Tyr
75                80                85

gcc aac cgc acg gcc ctc ttc ccg gac ctg ctg gca cag ggc aac gca      339
Ala Asn Arg Thr Ala Leu Phe Pro Asp Leu Leu Ala Gln Gly Asn Ala
90                95                100                105

tcc ctg agg ctg cag cgc gtg cgt gtg gcg gac gag ggc agc ttc acc      387
Ser Leu Arg Leu Gln Arg Val Arg Val Ala Asp Glu Gly Ser Phe Thr
110                115                120

tgc ttc gtg agc atc cgg gat ttc ggc agc gct gcc gtc agc ctg cag      435
Cys Phe Val Ser Ile Arg Asp Phe Gly Ser Ala Ala Val Ser Leu Gln
125                130                135

gtg gcc gct ccc tac tcg aag ccc agc atg acc ctg gag ccc aac aag      483
Val Ala Ala Pro Tyr Ser Lys Pro Ser Met Thr Leu Glu Pro Asn Lys
140                145                150

gac ctg cgg cca ggg gac acg gtg acc atc acg tgc tcc agc tac cag      531
Asp Leu Arg Pro Gly Asp Thr Val Thr Ile Thr Cys Ser Ser Tyr Gln
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ggc tac cct gag gct gag gtg ttc tgg cag gat ggg cag ggt gtg ccc      579
Gly Tyr Pro Glu Ala Glu Val Phe Trp Gln Asp Gly Gln Gly Val Pro
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ctg act ggc aac gtg acc acg tcg cag atg gcc aac gag cag ggc ttg      627
Leu Thr Gly Asn Val Thr Thr Ser Gln Met Ala Asn Glu Gln Gly Leu
190                195                200

ttt gat gtg cac agc atc ctg cgg gtg gtg ctg ggt gca aat ggc acc      675
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Gln Asp Gly Glu Gly Glu Gly Ser Lys Thr Ala Leu Gln Pro Leu Lys					
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<222> (422)..(422)

<223> The 'Xaa' at location 422 stands for Lys, Asn, Arg, Ser, Thr, Ile, Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon, Tyr, Trp, Cys, or Phe.

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<223> The 'Xaa' at location 423 stands for Lys, Asn, Arg, Ser, Thr, Ile, Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, a stop codon, Tyr, Trp, Cys, or Phe.

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<222> (424)..(424)

<223> The 'Xaa' at location 424 stands for Lys, Asn, Arg, Ser, Thr, Ile

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Tyr, Trp, Cys, or Phe.

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Tyr, Trp, Cys, or Phe.

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X11600.ST25.txt

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<220>

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<223> n = any nucleotide A, C, G, or T
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Val Pro Glu Asp Pro Val Val Ala Leu Val Gly Thr Asp Ala Thr Leu
35     40     45
Cys Cys Ser Phe Ser Pro Glu Pro Gly Phe Ser Leu Ala Gln Leu Asn

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 Leu Ile Trp Gln Leu Thr Asp Thr Lys Gln Leu Val His Ser Phe Ala  
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 Glu Gly Gln Asp Gln Gly Ser Ala Tyr Ala Asn Arg Thr Ala Leu Phe  
 85 90 95  
 Pro Asp Leu Leu Ala Gln Gly Asn Ala Ser Leu Arg Leu Gln Arg Val  
 100 105 110  
 Arg Val Ala Asp Glu Gly Ser Phe Thr Cys Phe Val Ser Ile Arg Asp  
 115 120 125  
 Phe Gly Ser Ala Ala Val Ser Leu Gln Val Ala Ala Pro Tyr Ser Lys  
 130 135 140  
 Pro Ser Met Thr Leu Glu Pro Asn Lys Asp Leu Arg Pro Gly Asp Thr  
 145 150 155 160  
 Val Thr Ile Thr Cys Ser Ser Tyr Gln Gly Tyr Pro Glu Ala Glu Val  
 165 170 175  
 Phe Trp Gln Asp Gly Gln Gly Val Pro Leu Thr Gly Asn Val Thr Thr  
 180 185 190  
 Ser Gln Met Ala Asn Glu Gln Gly Leu Phe Asp Val His Ser Ile Leu  
 195 200 205  
 Arg Val Val Leu Gly Ala Asn Gly Thr Tyr Ser Cys Leu Val Arg Asn  
 210 215 220  
 Pro Val Leu Gln Gln Asp Ala His Ser Ser Val Thr Ile Thr Pro Gln  
 225 230 235 240  
 Arg Ser Pro Thr Gly Ala Val Glu Val Gln Val Pro Glu Asp Pro Val  
 245 250 255  
 Val Ala Leu Val Gly Thr Asp Ala Thr Leu Arg Cys Ser Phe Ser Pro  
 260 265 270  
 Glu Pro Gly Phe Ser Leu Ala Gln Leu Asn Leu Ile Trp Gln Leu Thr  
 275 280 285  
 Asp Thr Lys Gln Leu Val His Ser Phe Thr Glu Gly Arg Asp Gln Gly  
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<213> Homo sapiens

<220>

<221> CDS

<222> (25)..(1629)

<223>

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<222> (25)..(1629)

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Met	Gly	Val	His	Val	Gly	Ala	Ala	Leu	Gly	Ala	Leu	
10				15				20			25	
aca	gga	gcc	ctg	gag	gtc	cag	gtc	cct	gaa	gac	cca	147
Thr	Gly	Ala	Leu	Glu	Val	Gln	Val	Pro	Glu	Asp	Pro	
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Val	Gly	Thr	Asp	Ala	Thr	Leu	Cys	Cys	Ser	Phe	Ser	
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ttc	agc	ctg	gca	cag	ctc	aac	ctc	atc	tgg	cag	ctg	243
Phe	Ser	Leu	Ala	Gln	Leu	Asn	Leu	Ile	Trp	Gln	Leu	
		60				65					70	
cag	ctg	gtg	cac	agc	ttt	gct	gag	ggc	cag	gac	cag	291
Gln	Leu	Val	His	Ser	Phe	Ala	Glu	Gly	Gln	Asp	Gln	
	75					80				85		
gcc	aac	cgc	acg	gcc	ctc	ttc	ccg	gac	ctg	ctg	gca	339
Ala	Asn	Arg	Thr	Ala	Leu	Phe	Pro	Asp	Leu	Leu	Ala	
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tcc	ctg	agg	ctg	cag	cgc	gtg	cgt	gtg	gcg	gac	gag	387
Ser	Leu	Arg	Leu	Gln	Arg	Val	Arg	Val	Ala	Asp	Glu	
			110						115			120
tgc	ttc	gtg	agc	atc	cgg	gat	ttc	ggc	agc	gct	gcc	435
Cys	Phe	Val	Ser	Ile	Arg	Asp	Phe	Gly	Ser	Ala	Ala	
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gtg	gcc	gct	ccc	tac	tcg	aag	ccc	agc	atg	acc	ctg	483
Val	Ala	Ala	Pro	Tyr	Ser	Lys	Pro	Ser	Met	Thr	Leu	
											Glu	
											Pro	
											Asn	
											Lys	

X11600.ST25.txt																
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gat Asp	ttc Phe	ggc Gly	agc Ser	gct Ala 350	gcc Ala	gtc Val	agc Ser	ctg Leu	cag Gln 355	gtg Val	gcc Ala	gct Ala	ccc Pro	tac Tyr 360	tcg Ser	1107
aag Lys	ccc Pro	agc Ser	atg Met 365	acc Thr	ctg Leu	gag Glu	ccc Pro	aac Asn 370	aag Lys	gac Asp	ctg Leu	cgg Arg	cca Pro 375	ggg Gly	gac Asp	1155
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gtg Val	ttc Phe	tgg Trp	cag Gln	gat Asp	ggg Gly	cag Gln	ggt Gly	gtg Val	ccc Pro	ctg Leu	act Thr	ggc Gly	aac Asn	gtg Val	acc Thr	1251

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Leu Arg Val Val Leu Gly Ala Asn Gly Thr Tyr Ser Cys Leu Val Arg					
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Asn Pro Val Leu Gln Gln Asp Ala His Gly Ser Val Thr Ile Thr Gly					
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Gln Pro Met Thr Phe Pro Pro Glu Ala Leu Trp Val Thr Val Gly Leu					
		460		465	470
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Ser Val Cys Leu Ile Ala Leu Leu Val Ala Leu Ala Phe Val Cys Trp					
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Arg Lys Ile Lys Gln Ser Cys Glu Glu Glu Asn Ala Gly Ala Glu Asp					
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Gln Asp Gly Glu Gly Glu Gly Ser Lys Thr Ala Leu Gln Pro Leu Lys					
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<220>

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<223>

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Val Pro Glu Asp Pro Val Val Ala Leu Val Gly Thr Asp Ala Thr Leu  
35 40 45

Cys Cys Ser Phe Ser Pro Glu Pro Gly Phe Ser Leu Ala Gln Leu Asn  
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Leu Ile Trp Gln Leu Thr Asp Thr Lys Gln Leu Val His Ser Phe Ala  
65 70 75 80

Glu Gly Gln Asp Gln Gly Ser Ala Tyr Ala Asn Arg Thr Ala Leu Phe  
85 90 95

Pro Asp Leu Leu Ala Gln Gly Asn Ala Ser Leu Arg Leu Gln Arg Val  
100 105 110

Arg Val Ala Asp Glu Gly Ser Phe Thr Cys Phe Val Ser Ile Arg Asp  
115 120 125

Phe Gly Ser Ala Ala Val Ser Leu Gln Val Ala Ala Pro Tyr Ser Lys  
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Pro Ser Met Thr Leu Glu Pro Asn Lys Asp Leu Arg Pro Gly Asp Thr  
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Val Thr Ile Thr Cys Ser Ser Tyr Gln Gly Tyr Pro Glu Ala Glu Val  
165 170 175

Phe Trp Gln Asp Gly Gln Gly Val Pro Leu Thr Gly Asn Val Thr Thr  
180 185 190

Ser Gln Met Ala Asn Glu Gln Gly Leu Phe Asp Val His Ser Ile Leu  
195 200 205

Arg Val Val Leu Gly Ala Asn Gly Thr Tyr Ser Cys Leu Val Arg Asn  
210 215 220

Pro Val Leu Gln Gln Asp Ala His Ser Ser Val Thr Ile Thr Pro Gln  
225 230 235 240

Arg Ser Pro Thr Gly Ala Val Glu Val Gln Val Pro Glu Asp Pro Val  
245 250 255

Val Ala Leu Val Gly Thr Asp Ala Thr Leu Arg Cys Ser Phe Ser Pro  
260 265 270

Glu Pro Gly Phe Ser Leu Ala Gln Leu Asn Leu Ile Trp Gln Leu Thr  
275 280 285

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Asp Thr Lys Gln Leu Val His Ser Phe Thr Glu Gly Arg Asp Gln Gly  
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 Ser Ala Tyr Ala Asn Arg Thr Ala Leu Phe Pro Asp Leu Leu Ala Gln  
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 Gly Asn Ala Ser Leu Arg Leu Gln Arg Val Arg Val Ala Asp Glu Gly  
 325 330 335  
 Ser Phe Thr Cys Phe Val Ser Ile Arg Asp Phe Gly Ser Ala Ala Val  
 340 345 350  
 Ser Leu Gln Val Ala Ala Pro Tyr Ser Lys Pro Ser Met Thr Leu Glu  
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 Pro Asn Lys Asp Leu Arg Pro Gly Asp Thr Val Thr Ile Thr Cys Ser  
 370 375 380  
 Ser Tyr Arg Gly Tyr Pro Glu Ala Glu Val Phe Trp Gln Asp Gly Gln  
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 Gly Val Pro Leu Thr Gly Asn Val Thr Thr Ser Gln Met Ala Asn Glu  
 405 410 415  
 Gln Gly Leu Phe Asp Val His Ser Val Leu Arg Val Val Leu Gly Ala  
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 Asn Gly Thr Tyr Ser Cys Leu Val Arg Asn Pro Val Leu Gln Gln Asp  
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 Ala His Gly Ser Val Thr Ile Thr Gly Gln Pro Met Thr Phe Pro Pro  
 450 455 460  
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 465 470 475 480  
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 485 490 495  
 Glu Glu Glu Asn Ala Gly Ala Glu Asp Gln Asp Gly Glu Gly Glu Gly  
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## X11600.ST25.txt

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gc	1682